

## REMARKS

By present amendment the claims have been amended to present the claims in accordance with customary U.S. practice, care having been exercised to avoid any introduction of new matter.

Claims 22-24 have been canceled, and Claims 1-21 and 25-27 have been amended as to form. Claims 28-38 have been added. Support for Claims 28 and 29 can be found in Claim 9, while support for Claims 30 and 31 can be found in Claim 11. Support for Claims 32-37 can be found in Claims 19, 20, 21, 25, 26, and 27, respectively. Support for Claim 38 can be found in the specification on page 6, lines 8-25. The amendments to the claims remove multiple dependencies and place the claims in United States form.

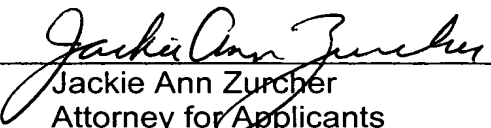
The specification has been amended to include section headings in accordance with customary U.S. practice, and to include a Summary of the Invention. Support for the amendment to the specification can be found in original Claim 1.

The amendments to the claims and the specification do not involve any introduction of new matter, whereby entry is believed to be in order and is respectively requested.

Attached is a marked-up version of the changes made to the specification and claims by the current amendment. The attached page is captioned "Version With Markings To Show Changes Made".

A disk containing a computer readable form of the Sequence Listing is enclosed herewith. The information recorded in computer readable form is identical to the Sequence Listing set forth on the paper copy. The attached diskette is submitted in American Standard Code for Information Interchange (ASCII) text.

Respectfully submitted,

By   
Jackie Ann Zurcher  
Attorney for Applicants  
Reg. No. 42,251

Bayer Corporation  
100 Bayer Road  
Pittsburgh, Pennsylvania 15205-9741  
(412) 777-8356  
FACSIMILE PHONE NUMBER:  
(412) 777-8363

/jme/JAZ0074

09847081-050204  
T03050 T3074860

VERSION WITH MARKINGS TO SHOW CHANGES MADE

**In the Claims:**

Claims 22-24 have been canceled.

Claims 1-21 and 25-27 have been amended as follows:

1. (Amended) A n[N]ucleic acid[s] which encodes a polypeptide[s] from tobacco with the bioactivity of a zeta-carotene desaturase, comprising [which comprises] the amino acid sequence of SEQ ID NO: 6.
2. (Amended) A n[N]ucleic acid[s] according to Claim 1, wherein the nucleic acid encodes a polypeptide [characterized in that they encode polypeptides] with the amino acid sequence of SEQ ID NO: 6.
3. (Amended) A n[N]ucleic acid[s] according to Claim 1, wherein the nucleic acid is a [1 or 2, characterized in that they are] single-stranded or double-stranded DNA or RNA.
4. (Amended) A n[N]ucleic acid[s] according to Claim 3, wherein the nucleic acid is a fragment [characterized in that they are fragments] of genomic DNA or cDNA.
5. (Amended) A n[N]ucleic acid[s] according to Claim 1, wherein the nucleic acid is [one of Claims 1 to 4, characterized in that they are] derived from tobacco plants.
6. (Amended) A n[N]ucleic acid[s] according to Claim 1 [one of Claims 1 to 5], comprising a sequence selected from [amongst]
  - (a) the sequence of SEQ ID NO: 5,

- (b) sequences encoding a polypeptide which comprises the amino acid sequence of SEQ ID NO: 6,
- (c) partial sequences of the sequences defined under (a) or (b) which are at least 14 base pairs in length,
- (d) sequences which hybridize with the sequences defined under (a), (b) or (c),
- (e) sequences which are complementary to the sequences defined under (a), (b) or (c), and
- (f) sequences which, owing to the degeneracy of the genetic code, encode the same amino acid sequence as the sequences defined under (a) to (c).

7. (Amended) A<sub>r</sub>[R]egulatory region which naturally controls, in plant cells, [in particular in tobacco plants, ]the transcription of a nucleic acid according to Claim 1 [one of Claims 1 to 6].

8. (Amended) A DNA construct comprising a nucleic acid according to Claim 1 [one of Claims 1 to 6 ]and a heterologous promoter.

9. (Amended) A<sub>v</sub>[V]ector comprising a nucleic acid according to Claim 1 [one of Claims 1 to 6, a regulatory region according to Claim 7 or a DNA construct according to Claim 8].

10. (Amended) A<sub>v</sub>[V]ector according to Claim 9, wherein [characterized in that] the nucleic acid is linked functionally to regulatory sequences which ensure the expression of the nucleic acid in pro- or eukaryotic cells.

11. (Amended) A<sub>h</sub>[H]ost cell containing a nucleic acid according to Claim 1 [one of Claims 1 to 6, a DNA construct according to Claim 8 or a vector according to Claim 9 or 10].

12. (Amended) A h[H]ost cell according to claim 11, wherein the host cell [characterized in that it] is a prokaryotic cell[, in particular *E. Coli*].

13. (Amended) A h[H]ost cell according to Claim 11, wherein the host cell [characterized in that it] is a eukaryotic cell[, in particular a yeast cell, insect cell, mammalian cell or plant cell].

14. (Amended) A p[P]olypeptide with the bioactivity of a phytoene synthase which is encoded by a nucleic acid of SEQ ID NO: 1 or SEQ ID NO: 3, comprising an amino acid sequence of SEQ ID NO: 2 or SEQ ID NO: 4.

15. (Amended) A p[P]olypeptide with the bioactivity of a zeta-carotene desaturase which is encoded by a nucleic acid according to Claim 1 [one of Claims 1 to 6].

16. (Amended) An a[A]ntibody which binds specifically to a polypeptide of Claim 14.

17. (Amended) An a[A]ntibody which binds specifically to a polypeptide of Claim 15.

18. (Amended) A p[P]rocess for generating a nucleic acid according to Claim 1, comprising the steps of [one of Claims 1 to 6, comprising the following steps]:

- (a) completely chemically synthesizing the nucleic acid [synthesis carried out in a manner known per se] or
- (b) chemically synthesizing [synthesis of] oligonucleotides, labelling the oligonucleotides, hybridizing the oligonucleotides with DNA of a genomic or cDNA library which has been generated starting from genomic DNA or mRNA

of plant cells, selecting [selection of] positive clones and isolating [isolation of] the hybridizing DNA from positive clones, or

- (c) chemically synthesizing [synthesis of] oligonucleotides and amplifying [amplification of the] target DNA by means of PCR.

19. (Amended) A p[P]rocess for generating a polypeptide according to Claim 14 [or 15], comprising

- (a1) culturing a host cell comprising a nucleic acid which encodes a polypeptide comprising the amino acid sequence of SEQ ID NO: 6 [according to one of Claims 11 to 13] under conditions which ensure the expression of the nucleic acid [according to one of Claims 1 to 6], or

- (a2[b]) expressing a nucleic acid which encodes a polypeptide comprising the amino acid sequence of SEQ ID NO: 6 [according to one of Claims 1 to 6] in an *in-vitro* system, and

- (b[c]) obtaining the polypeptide from the cell, the culture medium or the *in-vitro* system.

20. (Amended) A m[M]ethod of finding a chemical compound which binds to a polypeptide according to Claim 14 [and/or 15] or a polypeptide with the bioactivity of a phytoene desaturase, comprising the following steps

- (a) contacting a host cell containing a nucleic acid which encodes a polypeptide comprising the amino acid sequence of SEQ ID NO: 6 [according to one of Claims 11 to 13], a polypeptide according to Claim 14 [or 15] or a polypeptide with the bioactivity of a phytoene desaturase with a chemical compound or a mixture of chemical compounds under conditions which permit the interaction of a chemical compound with the polypeptide, and

- (b) determining the chemical compound which binds specifically to the polypeptide.

21. (Amended) A m[M]ethod of finding a compound which modifies the expression of polypeptides according to Claim 14 [or 15] or a polypeptide with the bioactivity of a phytoene desaturase, comprising the following steps:

- (a) contacting a host cell containing a nucleic acid which encodes a polypeptide comprising the amino acid sequence of SEQ ID NO: 6 [according to one of Claims 11 to 13] with a chemical compound or a mixture of chemical compounds,
- (b) determining the polypeptide concentration, and
- (c) determining the compound which specifically influences the expression of the polypeptide.

25. (Amended) An organism selected from t[T]ransgenic plants, parts of plants, protoplasts, plant tissues and [or] plant propagation materials, wherein the organism comprises an [characterized in that the] intracellular concentration of a polypeptide according to Claim 16 [or 17] which is increased or reduced in comparison with the corresponding wild-type cells after introducing a nucleic acid which encodes a polypeptide comprising the amino acid sequence of SEQ ID NO: 6 [according to one of Claims 1 to 6, a DNA construct according to Claim 8 or a vector according to Claim 9].

26. (Amended) An organism selected from p[P]lants, parts of plants, protoplasts, plant tissues or plant propagation materials, wherein the organism comprises [characterized in that they contain] a polypeptide according to Claim 14 [or 15] whose bioactivity or expression pattern is modified in comparison with the corresponding endogenous polypeptides.

27. (Amended) A method of generating plants, parts of plants, protoplasts, plant tissues or plant propagation materials according to Claim 25, comprising the step of modifying a nucleic acid which encodes a polypeptide comprising the amino acid sequence of SEQ ID NO: 6 by endogenous mutagenesis [characterized in that a nucleic acid according to one of Claims 1 to 6 or a regulatory region according to Claim 7 is modified by endogenous mutagenesis].

Claims 28-38 are new and have been added as follows:

--28. A vector comprising a regulatory region according to Claim 7.

29. A vector comprising a DNA construct according to Claim 8.

30. A host cell containing a DNA construct according to Claim 8.

31. A host cell containing a vector according to Claim 9.

32. A process for generating a polypeptide according to Claim 15, comprising

(a1) culturing a host cell comprising a nucleic acid comprising a sequence selected from:

- (i) the sequence of SEQ ID NO: 5,
- (ii) sequences encoding a polypeptide which comprises the amino acid sequence of SEQ ID NO: 6,
- (iii) sequences which hybridize with the sequences defined under (i) or (ii),
- (iv) sequences which are complementary to the sequences defined under (i) or (ii), and
- (v) sequences which, owing to the degeneracy of the genetic code, encode the same amino acid sequence as the sequences defined under (i) or (ii);

in a culture medium under conditions which ensure the expression of the nucleic acid, or

Mo6314



- (a2) expressing a nucleic acid which encodes a polypeptide with the bioactivity of a zeta-carotene desaturase, comprising the amino acid sequence of SEQ ID NO: 6 in an *in-vitro* system, and
- (b) obtaining the polypeptide from the cell, the culture medium or the *in-vitro* system.

33. A method of finding a chemical compound which binds to a polypeptide according to Claim 15 or a polypeptide with the bioactivity of a phytoene desaturase, comprising the following steps:

- (a) contacting a host cell comprising a nucleic acid which encodes a polypeptide with the bioactivity of a zeta-carotene desaturase, comprising the amino acid sequence of SEQ ID NO: 6, a polypeptide according to Claim 15 or a polypeptide with the bioactivity of a phytoene desaturase with a chemical compound or a mixture of chemical compounds under conditions which permit the interaction of a chemical compound with the polypeptide, and
- (b) determining the chemical compound which binds specifically to the polypeptide.

34. A method of finding a compound which modifies the expression of polypeptides according to Claim 15 or a polypeptide with the bioactivity of a phytoene desaturase, comprising the following steps:

- (a) contacting a host cell containing a nucleic acid which encodes a polypeptide comprising the amino acid sequence of SEQ ID NO: with a chemical compound or a mixture of chemical compounds,
- (b) determining the polypeptide concentration, and

- (c) determining the compound which specifically influences the expression of the polypeptide.

35. An organism selected from transgenic plants, parts of plants, protoplasts, plant tissues and plant propagation materials, wherein the organism comprises an intracellular concentration of a polypeptide according to Claim 17 which is increased or reduced in comparison with the corresponding wild-type cells after introducing a nucleic acid which encodes a polypeptide comprising the amino acid sequence of SEQ ID NO: 6.

36. An organism selected from plants, parts of plants, protoplasts, plant tissues or plant propagation materials, wherein the organism comprises a polypeptide according to Claim 15 whose bioactivity or expression pattern is modified in comparison with the corresponding endogenous polypeptides.

37. A method of generating plants, parts of plants, protoplasts, plant tissues or plant propagation materials according to Claim 25, comprising the step of modifying by endogenous mutagenesis a regulatory region which naturally controls, in plant cells, the transcription of a nucleic acid which encodes a polypeptide comprising the amino acid sequence of SEQ ID NO: 6.

38. A nucleic acid comprising a sequence selected from the group consisting of:

- (a) the sequences of SEQ ID NOS: 1, 3 and 5,
- (b) sequences encoding polypeptides which comprises the amino acid sequences of SEQ ID NOS: 2, 4, and 6, and
- (c) sequences which, owing to the degeneracy of the genetic code, encode the same amino acid sequence as the amino acid sequences of SEQ ID NOS: 2, 4, and 6.